

8757-007

00/738944

SH 1 / OF 22

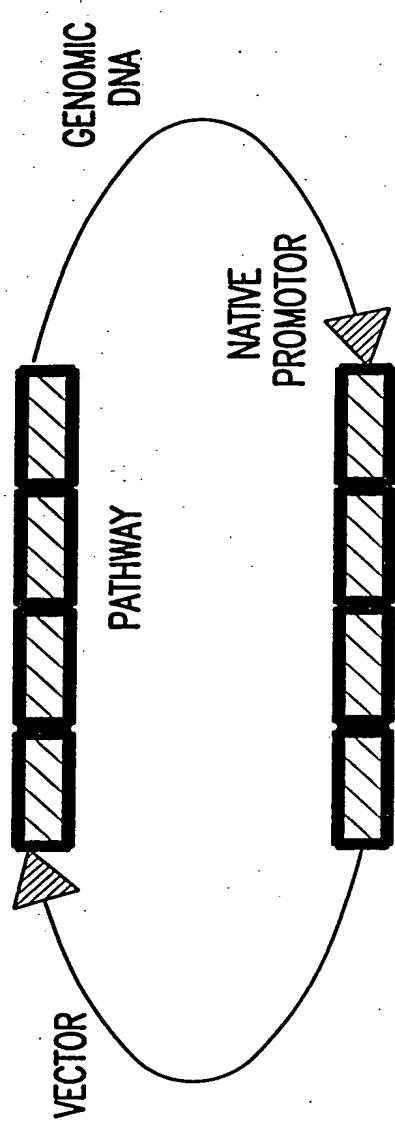


FIG. 1

26

8757-007

08/738944
SHEET 2 OF 22

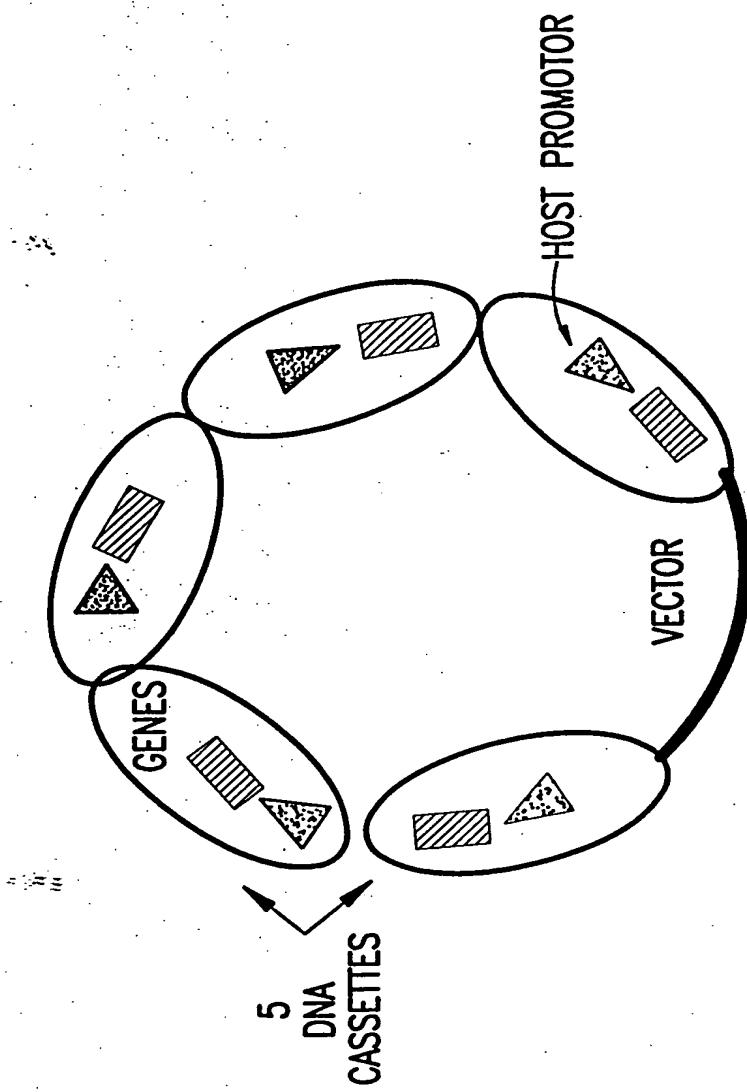


FIG. 2

8757-007

18/738944
SHEET 3 OF 22

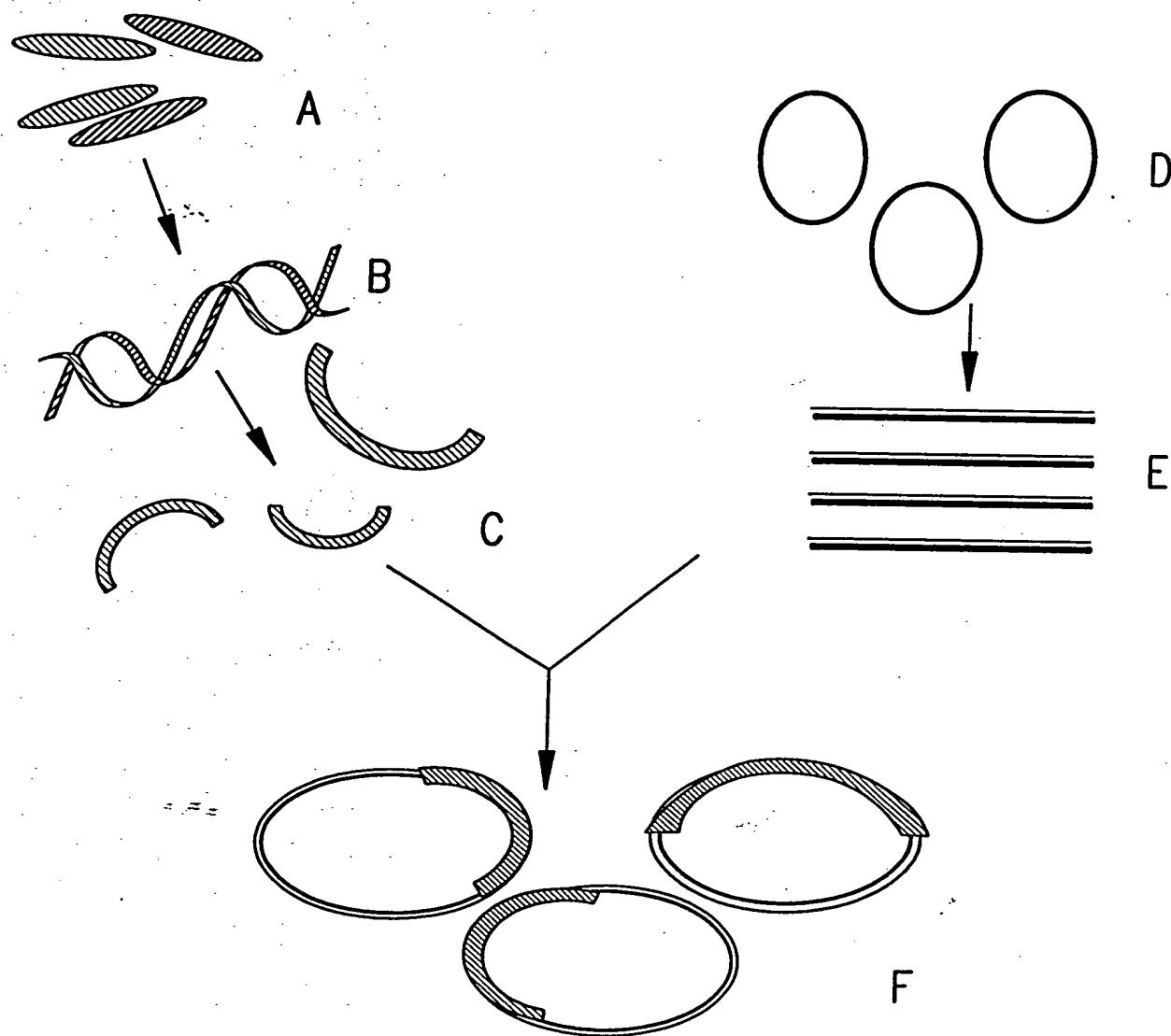


FIG.3

8757-007

08/738944

SHEET 4 OF 22

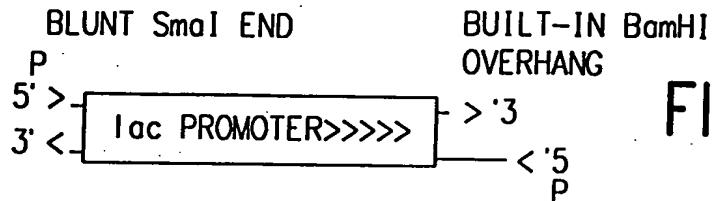


FIG. 4A

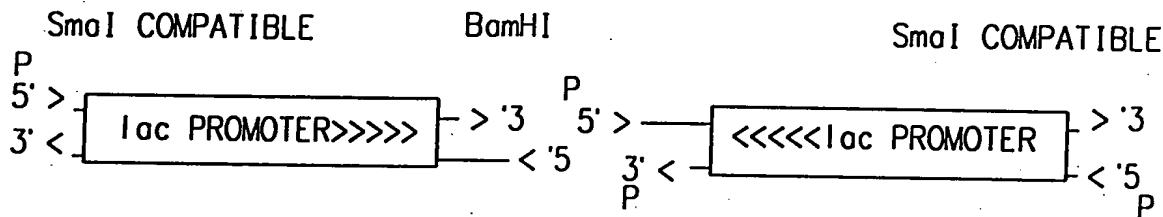


FIG.4B

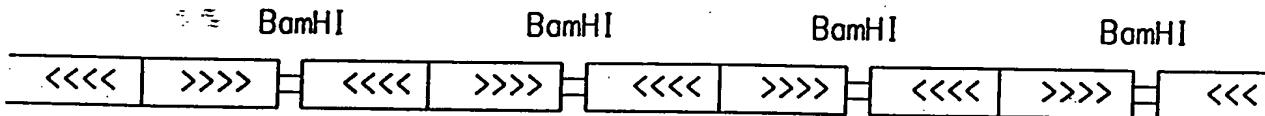


FIG. 4C

8757-007

08/738944

SHEET 5 OF 22

PROMOTERS FOR
cDNA & gDNA INSERTS

5' - GAGTAGAIC | PCR PROMOTER .. CTCGAGCGG-3'
3' - CTCACTAGA | ... FRAGMENT ... GAGCTGGCG-5'

Xba I

CUT w/Bgl II & Xba I

P
5' - GATCT | PCR PROMOTER .. CTC-3'
3' - A | ... FRAGMENT ... GAGCT-5'
Bgl II

Xba I
FILL IN WITH dTTP & dCTP

P
5' - GATCT | PCR PROMOTER .. CTC-3'
3' - A | ... FRAGMENT ... GAGCT-5'
Bgl II

Xba I
TREAT w/ PHOSPHATASE

5' - GATCT | PCR PROMOTER .. CTC-3'
3' - A | ... FRAGMENT ... GAGCT-5'
Bgl II

Xba I

PROMOTORS READY TO
LIGATE TO INSERTS

DISSIMILAR ENZYMES ON PROMOTER &
TERMINATOR FRAGMENTS ASSURE
DIRECTIONAL CLONING OF cDNA INSERTS.
(FOR EXAMPLE Xba I & Xba I)

ENZYME CLEAVAGE GENERATES
DEFINED ENDS, LEAVING
PROTECTED 3' BamHI SITE

KLENOW FILL IN OF
PROMOTERS & TERMINATORS
FRAGMENTS MAKE THEM INCAPABLE
OF INTER/INTRA LIGATION

PHOSPHATASE TREATMENT
CRATES EQUAL STRENGTH
LIGATION PARTNERS

5' - GATCCCCGGG | PCR TERMINATOR .. GGATCCCGGG-3'
3' - CTAGGGCCC | ... FRAGMENT ... CCTAGGGCCG-5'

BamHI

Xba I
CUT ONLY WITH Xba I

P
5' - CCCGG | PCR TERMINATOR .. GGATCCCGGG-3'
3' - C | ... FRAGMENT ... CCTAGGGCCG-5'

BamHI

Xba I
FILL IN WITH dCTP

P
5' - CCCGG | PCR TERMINATOR .. GGATCCCGGG-3'
3' - CCC | ... FRAGMENT ... CCTAGGGCCG-5'

BamHI

Xba I
TREAT w/ PHOSPHATASE

BamHI

TERMINATORS READY TO
LIGATE TO cDNA INSERTS

FIG.5A

8757-007

08/738944

SHEET 6 OF 22

TERMINATORS

5' - CAGTACATCT . PCR PROMOTER . CTCAGGGC-3'
 3' - CTCATACTAGA ... FRAGMENT ... GAGCTCGCG-5'

Bgl II Xho I

CUT W/Bgl II & Xho I

P
5' - GATCT . PCR PROMOTER . CTC-3'.
3' - A... FRAGMENT ... GAGCT-5'.
Bgl II Xho I

FILL IN WITH dTTP & dCTP

ENZYME CLEAVAGE GENERATES
DEFINED ENDS, LEAVING
PROTECTED 3' BamHI SITE

P
5' - GATCT . PCR PROMOTER . CTC-3'.
3' - A... FRAGMENT ... GAGCT-5'.
Bgl II Xho I

TREAT W/ PHOSPHATASE

KLENOW FILL IN OF
PROMOTERS & TERMINATORS
FRAGMENTS MAKE THEM INCAPABLE
OF INTER/INTRA LIGATION

P
5' - GATCT . PCR PROMOTER . CTC-3'.
3' - A... FRAGMENT ... GAGCT-5'.
Bgl II Xho I

PHOSPHATASE TREATMENT
CREATES EQUAL STRENGTH
LIGATION PARTNERS

5' - CGCCCTCGAG . PCR TERMINATOR . GGATCCGGG-3'
3' - CCCGAGCT . FRAGMENT ... CCTAGGGCGG-5'

BamHI Xho I

CUT ONLY WITH Xho I

P
5' - TCCAG . PCR TERMINATOR . GGATCCGGG-3'.
3' - C... FRAGMENT ... CCTAGGGCGG-5'.
Xho I BamHI P

FILL IN WITH dTTP & dCTP

P
5' - TCGAG . PCR TERMINATOR . GGATCCGGG-3'.
3' - CTC . FRAGMENT ... CCTAGGGCGG-5'.
Xho I BamHI P

P
5' - TCGAG . PCR TERMINATOR . GGATCCGGG-3'.
3' - CTC . FRAGMENT ... CCTAGGGCGG-5'.
Xho I BamHI

TREAT W/ PHOSPHATASE

PROMOTORS READY TO
LIGATE TO INSERTS

FIG. 5B

TERMINATORS READY TO
LIGATE TO INSERTS

8757-007

SHEET 7 OF 22

08/738944

FIRST STRAND cDNA SYNTHESIS PREPARED WITH Not I CONTAINING poly-dT PRIMER AND 5'-dCTP, AFTER 2nd STRAND SYNTHESIS, MODIFIED BamHI ADAPTERS ARE ADDED & cDNA IS DIGESTED WITH Not I, GIVING DIRECTIONAL cDNA GENE INSERTS

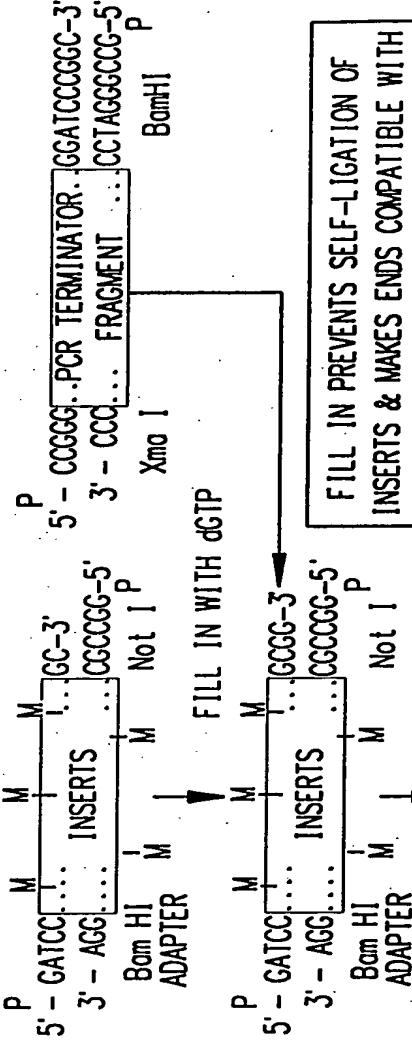
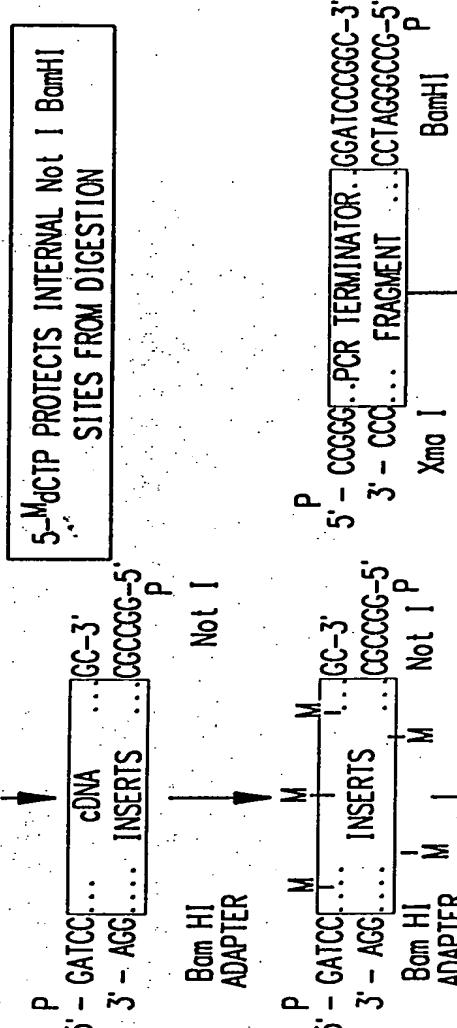
INSERT cDNAs

5' - GATC**T** PCR PROMOTER **CTC-3'**
3' - A... FRAGMENTS ... GAGCT-5'
Bgl II

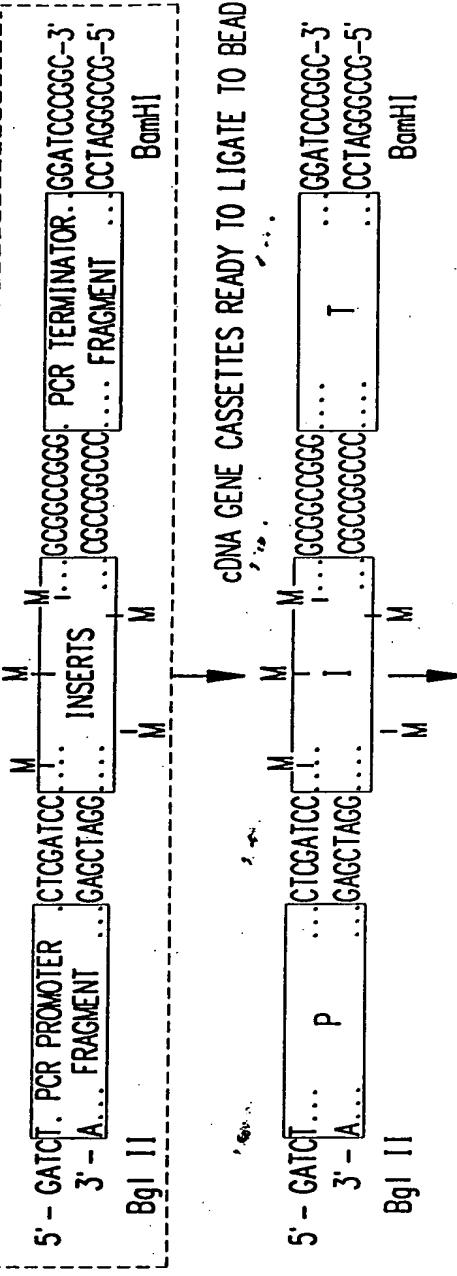
5' - GATC**C** PCR PROMOTER **GAGCT-3'**
3' - A... FRAGMENTS ... CCTAG-5'
Bgl II

FIG. 5C

5'-MdCTP PROTECTS INTERNAL Not I BamHI SITES FROM DIGESTION



FILL IN PREVENTS SELF-LIGATION OF INSERTS & MAKES ENDS COMPATIBLE WITH DIRECTIONAL PRO & TER PCR FRAGMENTS



SAMPLE cDNA GENE CASSETTES

5' - GATC**T** PCR PROMOTER **CTC-3'**
3' - A... FRAGMENT ... CCTAG-5'
Bgl II

5' - GATC**C** PCR PROMOTER **GAGCT-3'**
3' - A... FRAGMENT ... GAGCTAGG...
Bgl II



CDNA GENE CASSETTES READY TO LIGATE TO BEAD

5' - GATC... P
5' - GATC... M... CGCGCG... 1
3' - A... GAGCTAGG... 1
Bgl II

5' - GATC... P
5' - GATC... M... CGCGCG... 1
3' - A... GAGCTAGG... 1
Bgl II

BamHI

BamHI

BamHI

BamHI

08/738944

8757-007

SHEET 8 OF 22

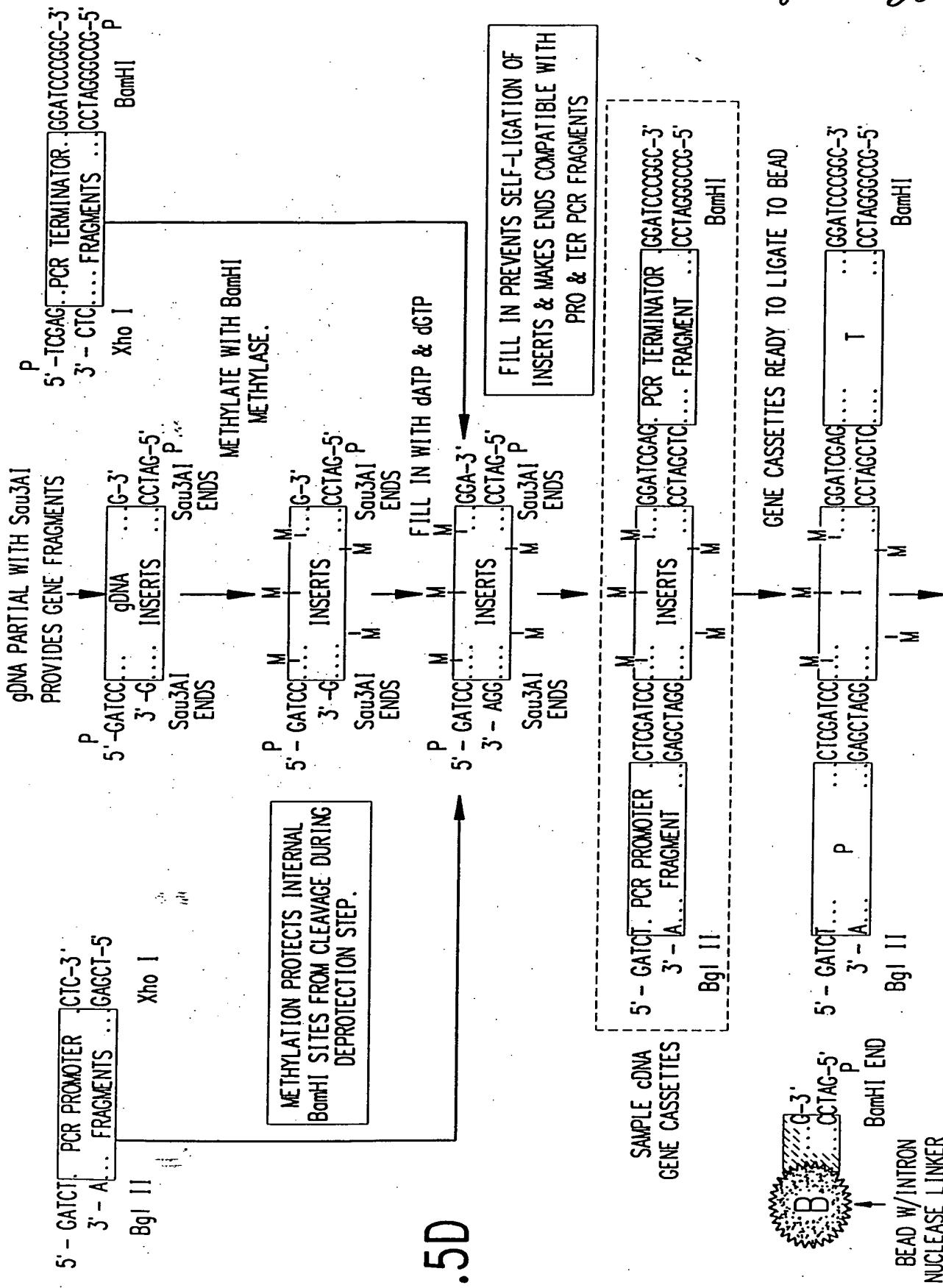


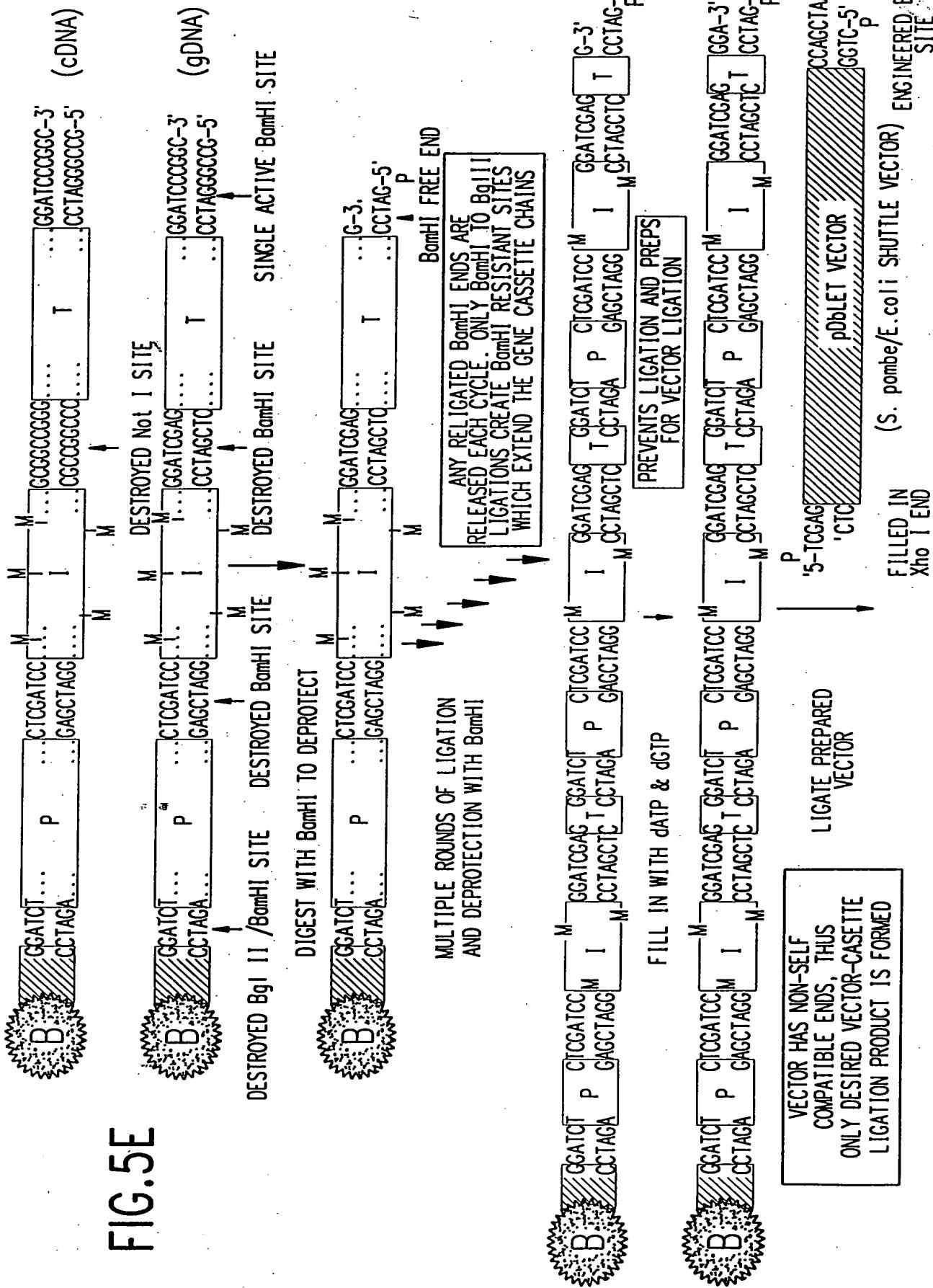
FIG. 5D

Due: 1

08/738944

8757-007

SHEET 9 OF 22

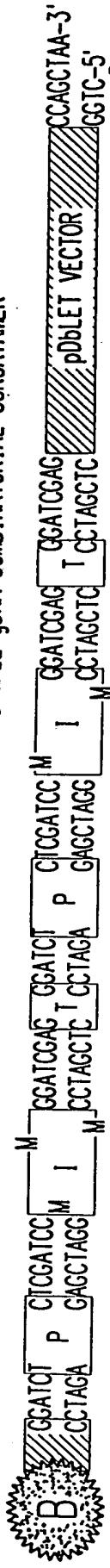


08/738944

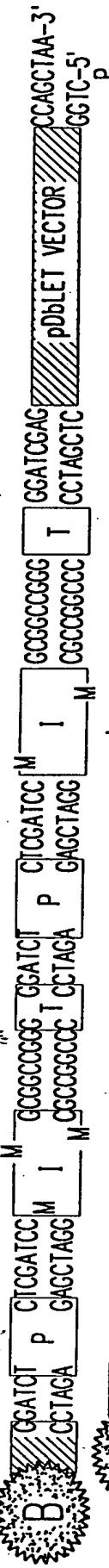
8757-007

SHEET 10 OF 22

SAMPLE gDNA COMBINATORIAL CONCATAMER



SAMPLE cDNA COMBINATORIAL CONCATAMER



↓
CUT WITH INTRON NUCLEASE TO RELEASE
SOLID-PHASE LIGATION BEAD
(SINGLE SITE IN LINKER OLIGO)

→ D GCTAA-3'
G-5'

SPIN AWAY & REMOVED

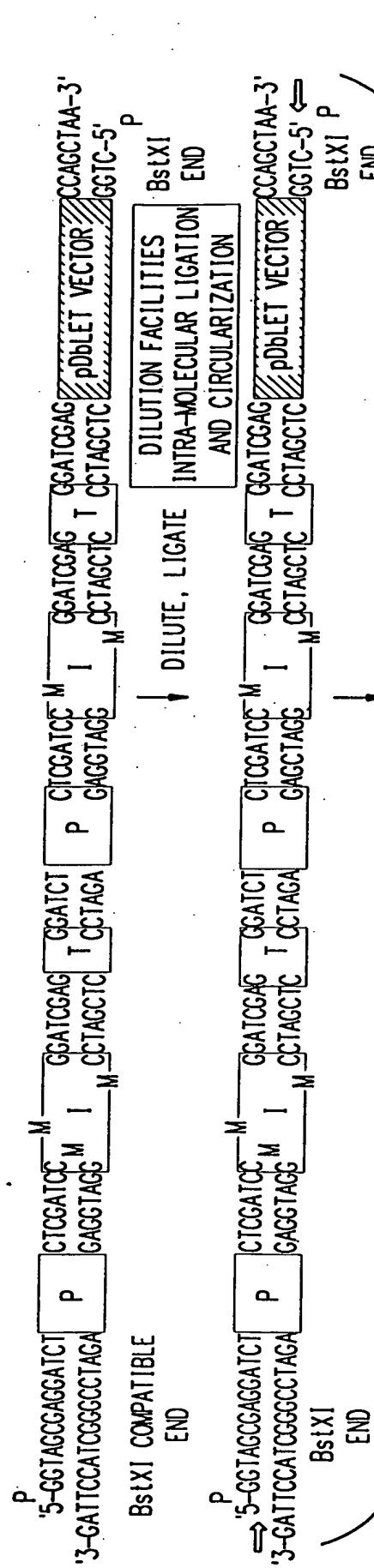


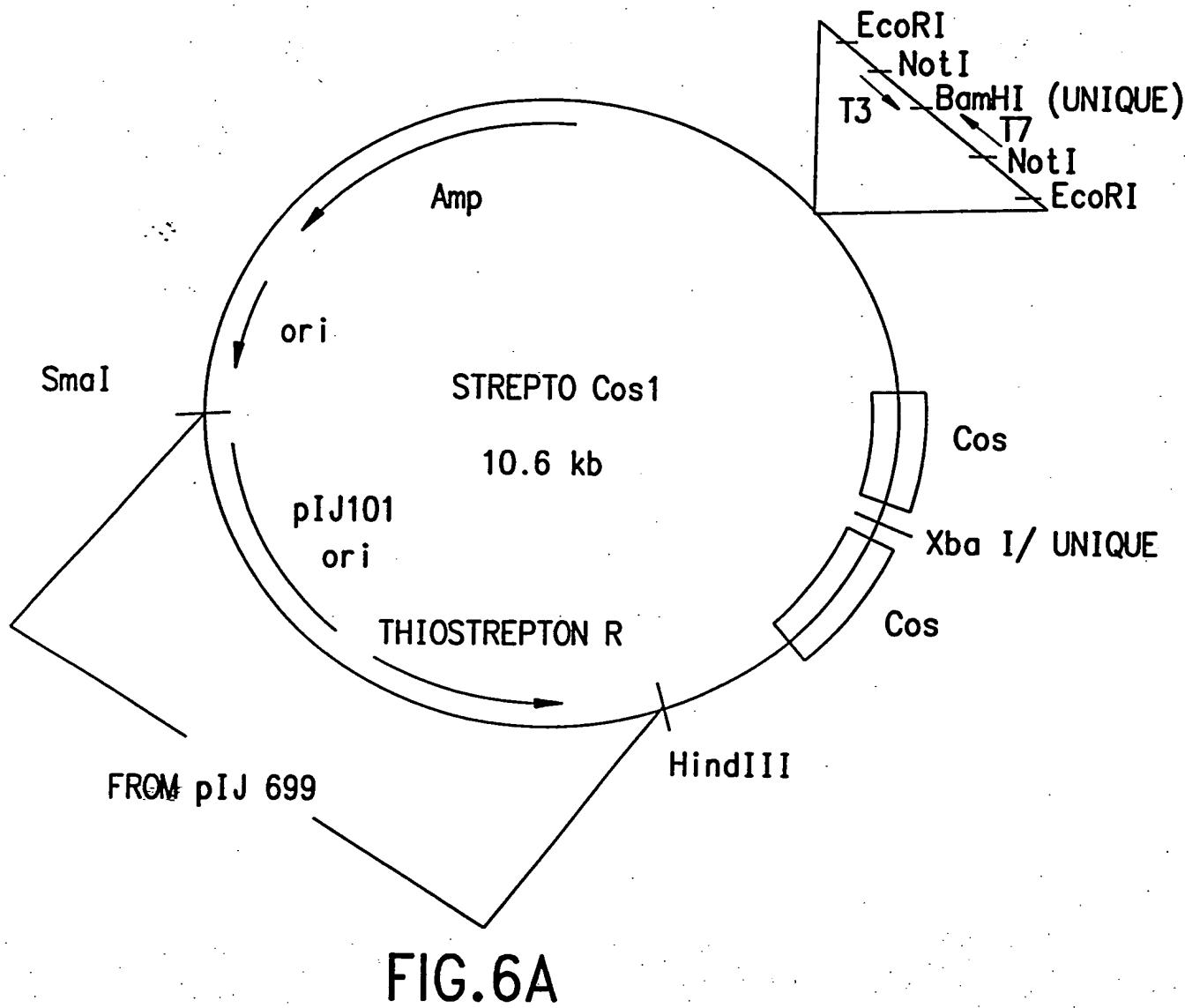
FIG. 5F

TRANSFORM *S. pombe* AND/OR *E. coli* AND
SCREEN RESULTING CLONES FOR
COMBINATORIAL ACTIVITIES

08/738944

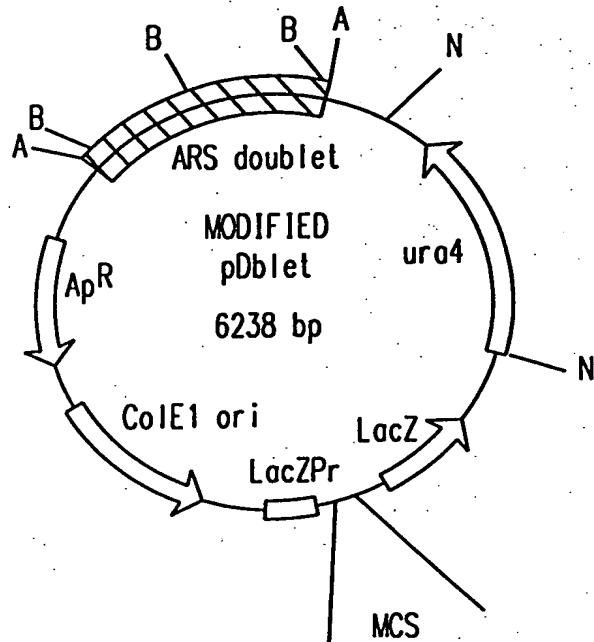
8757-007

SHEET 11 OF 22



8757-007

08/738944
SHEET 12 OF 22



MCS = SacI-NcoI-BstXI-NotI-Xba...

FIG.6B

5' CCTAGCCATGGCCACCTAACTGGGATGCC 3'
3' TCGAGGATCGGTACCGTGGATTGACCCTAGCGCCGG 5'
SacI NcoI BstXI NotI END

FIG.6C

8757-007

11/738944
SHEET 13 OF 22

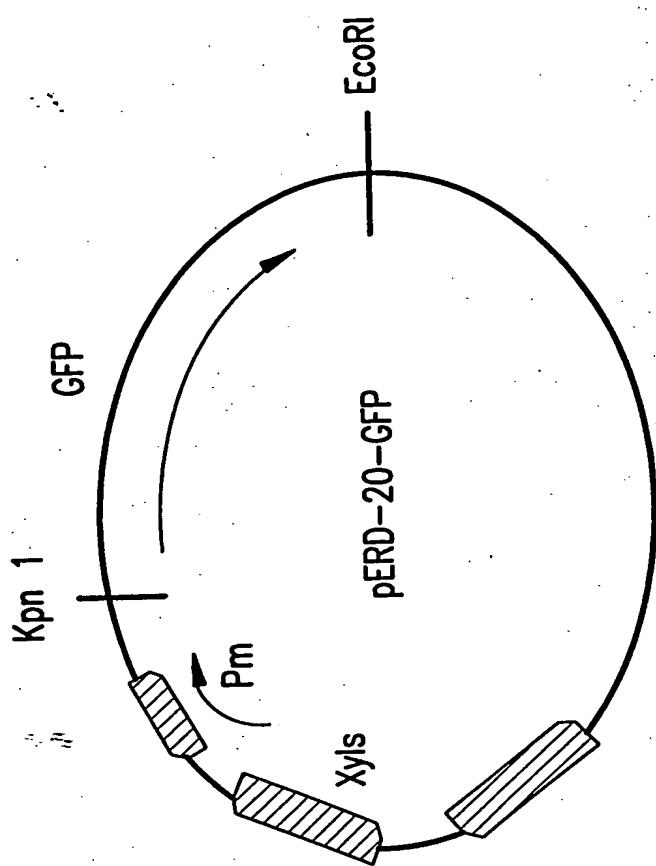


FIG. 7

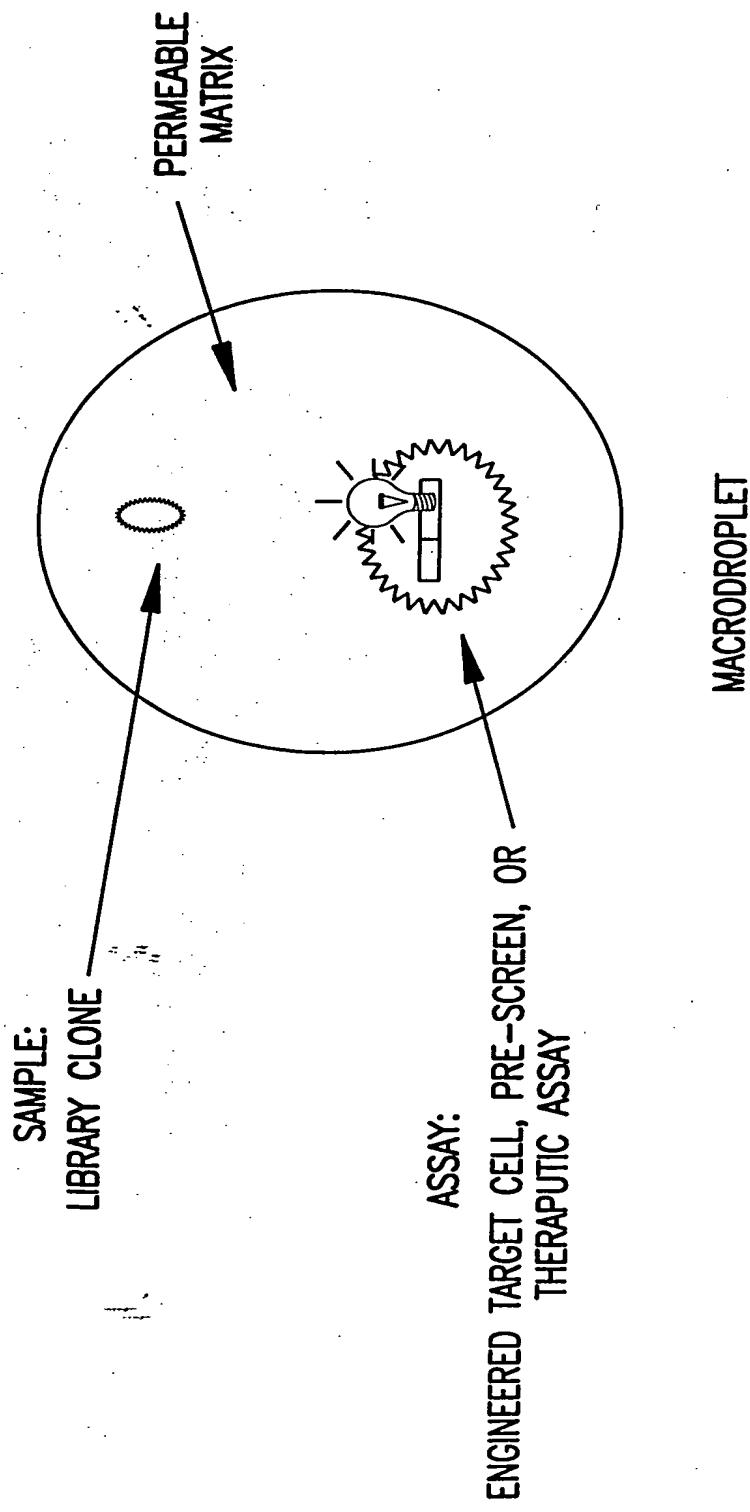


FIG. 8

08/738944

8757-007

SHEET 15 OF 22

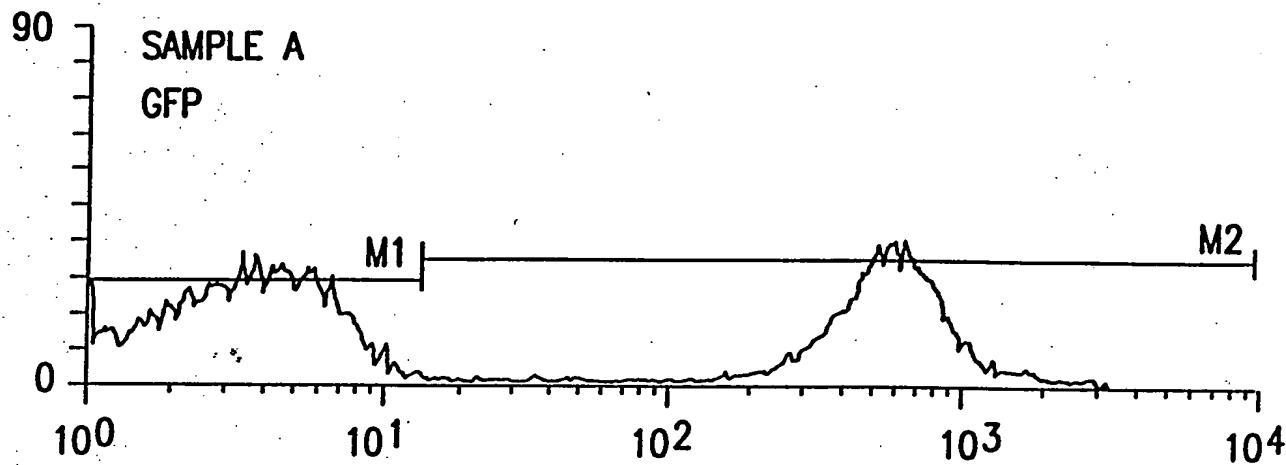


FIG.9A

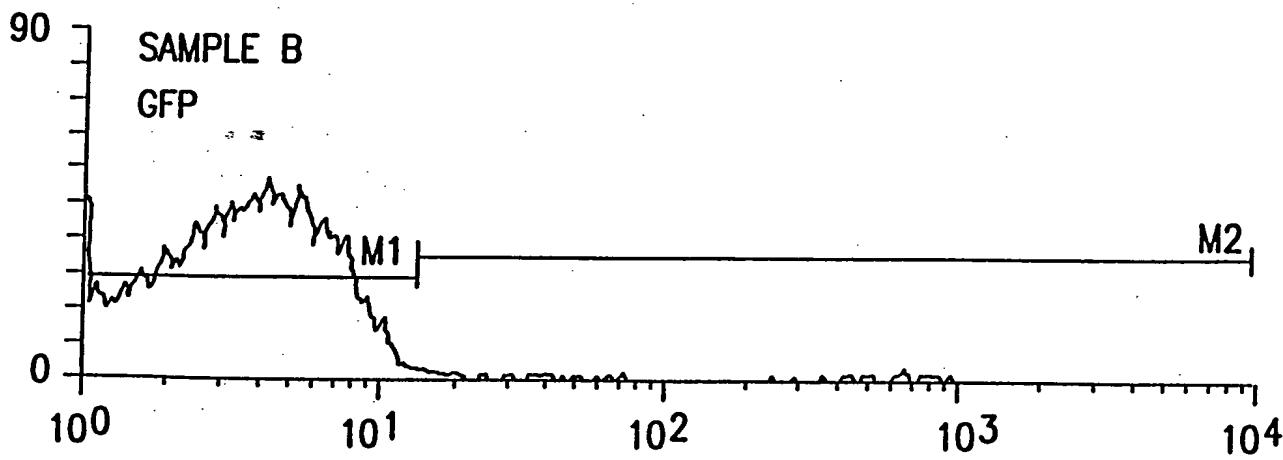


FIG.9B

08/738944

8757-007

SHEET 17 OF 22

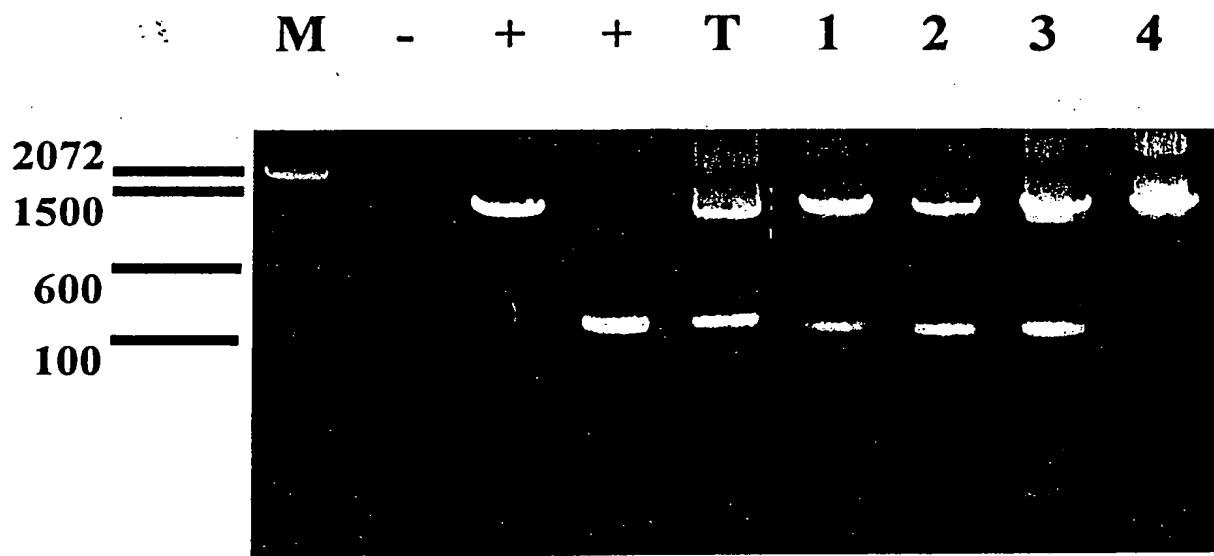


FIG.11

8757-007

SHEET 18 OF 22 08/738944

POOL 1

M - + + 1 2 3 4 5 6 7 8 9 10



2072
1500
600
100

FIG. 12A

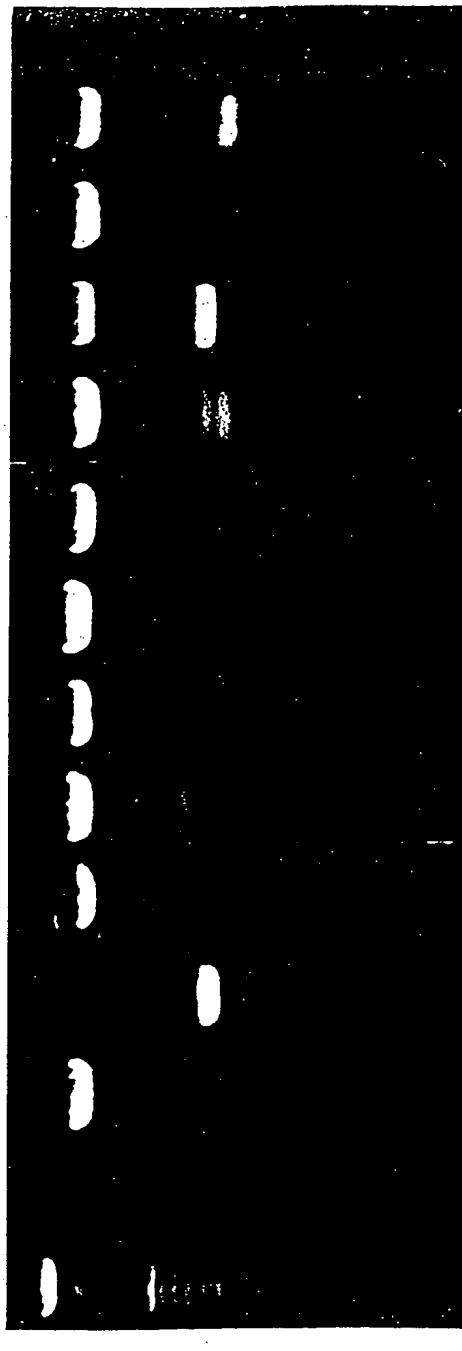
10/738944

8757-007

SHEET 19 OF 22

POOL 2

M - + + 12 13 14 15 16 17 18 19 20



2072
1500
600
100

FIG. 12B

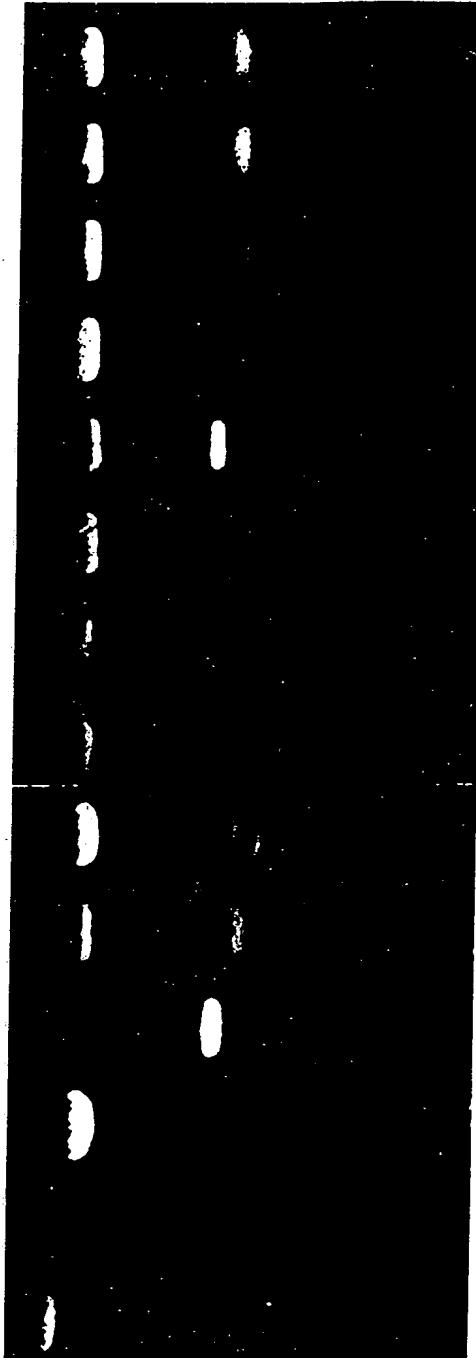
08/738944

8757-007

SHEET 20 OF 22

POOL 3

M - + + 21 22 23 27 30 31 32 33 34 35



2072
1500
600
100

FIG. 12C

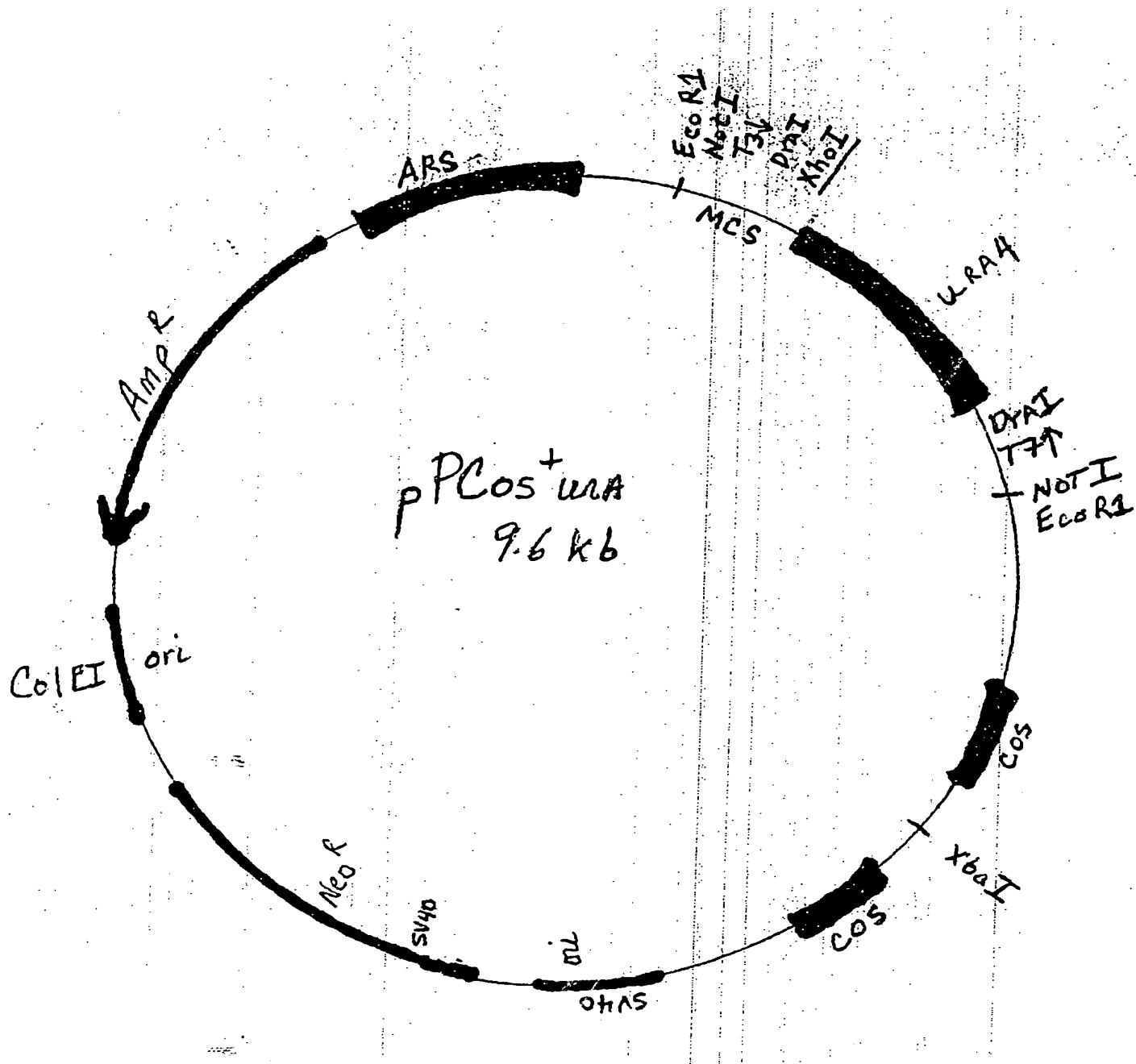


FIG. 13

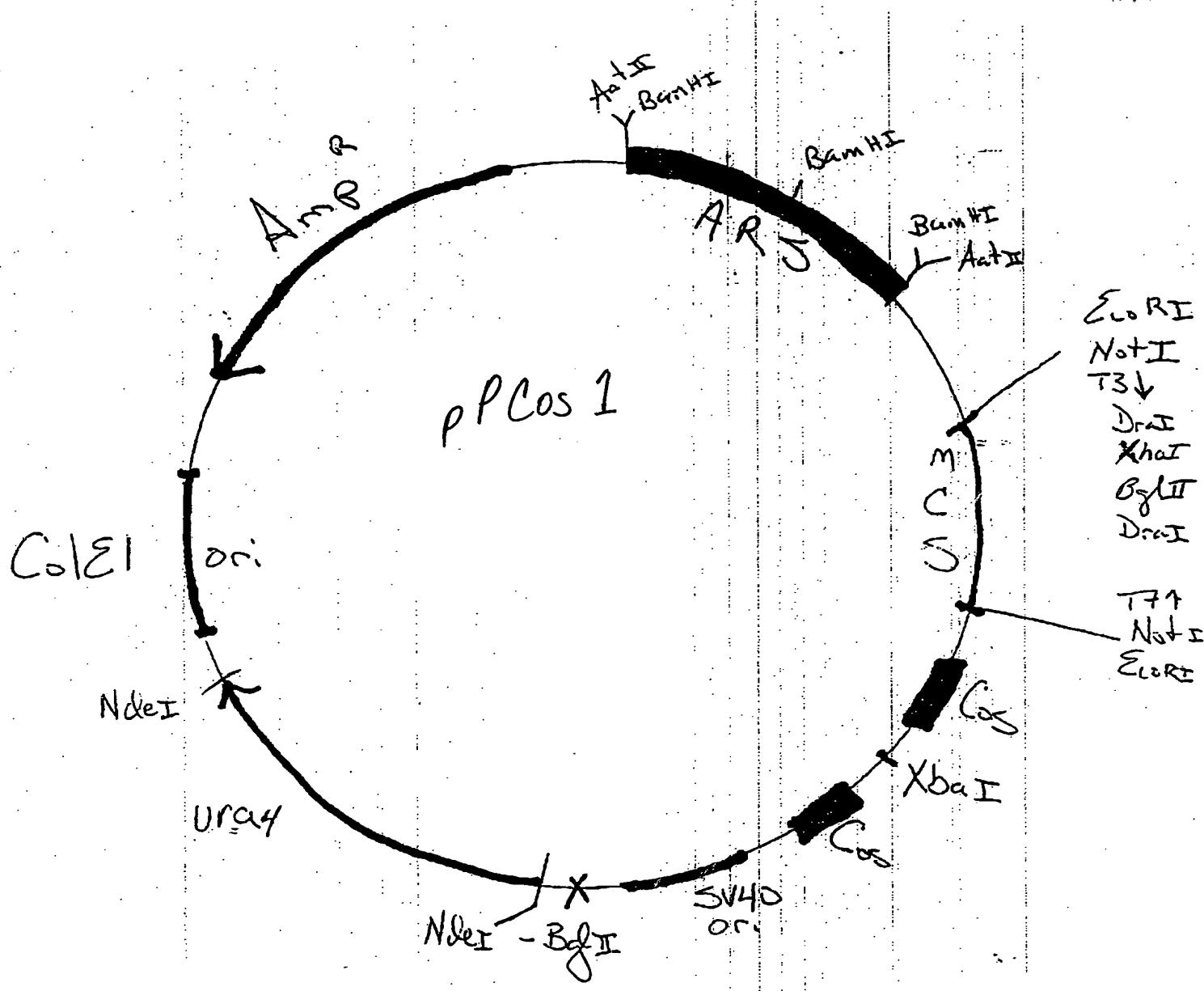


FIG. 14